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Alignment View Print

seq1	1 10 20 30
seq25	C C T T C T C C	G G C C C T T C C C	C T C G G C T T T G
Consensus	A A A A A A A A A A	A A A A A A A A A A	A A A A A A A A A A
seq1 40 50 60
seq25	C C T G G C G C	T C C T G C C T C C	C G C G G G C C C
Consensus	A A A A A A A A A A	A A A A A A A A A A	A A A A A A A A A A
seq1 70 80 90
seq25	C C T G T G T C C	C C C G C C C C C	C T C C C C C G
Consensus	A A A A A A A A A A	A A A A A A A A A A	A A A A A A A A A A
seq1 100 110 120
seq25	C G G C C T G G	C C C C T C T C T G	C T G C C G G C
Consensus	A A A A A A A A A A	A A A A A A A A A A	A A A A A A A A A A
seq1 130 140 150
seq25	C C C C C T G C T G	C C C C T C T C C	T G C T G C T C G G
Consensus	A A A A A A A A A A	A A A A A A A A A A	A A A A A A A A A A
seq1 160 170 180
seq25	G T T C T G G G C	C G C C T T G T C	C C C G G G C
Consensus	A A A A A A A A A A	A A A A A A A A A A	A A A A A A A A A A
seq1 190 200 210
seq25	G G T T C T C T T	T C T C T T T C T C	T T C T G G C C C
Consensus	A A A A A A A A A A	A A A A A A A A A A	A A A A A A A A A A

Alignment View Print

seq1 seq25	<div> <div>.....220</div> <div>C G C C C C C</div> </div>	<div> <div>.....230</div> <div>T G C C C T G</div> </div>	<div> <div>.....240</div> <div>G T T C C T C T</div> </div>
Consensus	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>
seq1 seq25	<div> <div>.....250</div> <div>C T G G G T T C</div> </div>	<div> <div>.....260</div> <div>T C C T G C T G C</div> </div>	<div> <div>.....270</div> <div>T G G C T T C C C G</div> </div>
Consensus	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>
seq1 seq25	<div> <div>.....280</div> <div>G C T G C C C G G T</div> </div>	<div> <div>.....290</div> <div>C T G G C C T G</div> </div>	<div> <div>.....300</div> <div>G G C C T G G G C</div> </div>
Consensus	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>
seq1 seq25	<div> <div>.....310</div> <div>C T C C T G G G</div> </div>	<div> <div>.....320</div> <div>C T G G G G G C C T</div> </div>	<div> <div>.....330</div> <div>C C T C C C T G G</div> </div>
Consensus	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>
seq1 seq25	<div> <div>.....340</div> <div>G T C C C C C C</div> </div>	<div> <div>.....350</div> <div>C G C C G C</div> </div>	<div> <div>.....360</div> <div>G T C T T G G C</div> </div>
Consensus	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>
seq1 seq25	<div> <div>.....370</div> <div>T G G T G C T G T</div> </div>	<div> <div>.....380</div> <div>T C T C C C T T C</div> </div>	<div> <div>.....390</div> <div>C T G G G G C C C</div> </div>
Consensus	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>
seq1 seq25	<div> <div>.....400</div> <div>C C T G C T C G C</div> </div>	<div> <div>.....410</div> <div>C C C G C T C T C</div> </div>	<div> <div>.....420</div> <div>C C G T C C T G C</div> </div>
Consensus	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>	<div> <div>.....</div> <div>.....</div> </div>



Alignment View Print

	430	440	450
seq1			
seq25			
Consensus			
seq1			
seq25			
Consensus			
seq1			
seq25			
Consensus			
seq1			
seq25			
Consensus			
seq1			
seq25			
Consensus			
seq1			
seq25			
Consensus			
seq1			
seq25			
Consensus			

Alignment View Print

	640	650	660
seq1	C C C C C C	C C C C T C T C T	C C C C C C C C
seq25	C C C C C C	C C C C T C T C T	C C T T C C C C
Consensus	C C C C C C	C C C C T C T C T	C C T C C C C C
	670	680	690
seq1	C C C C C C	C T C C T C C T C	C C
seq25	C C C C C C	C T C C T C C T C	C C
Consensus	C C C C C C	C T C C T C C T C	C C
	700	710	720
seq1	C C C C C C	C C C C T C C C C	C C T T C C C C
seq25	C C C C C C	C C C C T C C C C	C C T T C C C C
Consensus	C C C C C C	C C C C T C C C C	C C T T C C C C
	730	740	750
seq1	C C T C C C C C	C C C C C C C C	C T C C C C C C
seq25	C C T C C C C C	C C C C C C C C	C T C C C C C C
Consensus	C C T C C C C C	C C C C C C C C	C T C C C C C C
	760	770	780
seq1	C C C C C C C C	C C C C C C C C	C C C T C T C T C
seq25	C C C C C C C C	C C C C C C C C	C C C T C T C T C
Consensus	C C C C C C C C	C C C C C C C C	C C C T C T C T C
	790	800	810
seq1	C C C C C C C C	C C C C C C C C	T T C T C T C C C
seq25	C C C C C C C C	C C C C C C C C	T T C T C T C C C
Consensus	C C C C C C C C	C C C C C C C C	T T C T C T C C C
	820	830	840
seq1	T C C C C C C T	C C C C C C C T	C C C C C C C
seq25	T C C C C C C T	C C C C C C C T	C C C C C C C
Consensus	T C C C C C C T	C C C C C C C T	C C C C C C C



Alignment View Print

	850	860	870
seq1	C C T C C	C T C C T C	C C C C C
seq25	C C T C C	C T C C T C	C C C C C
Consensus	C C T C C	C T C C T C	C C C C C
	880	890	900
seq1	C C T C C C	C C C C T C C	C C T C C T T
seq25	C C T C C C	C C C C T C C	C C T C C T T
Consensus	C C T C C C	C C C C T C C	C C T C C T T
	910	920	930
seq1	C C T C C T T T C T	C T C C C C C C	C T C C T C T C C
seq25	C C T C C T T T C T	C T C C C C C C	C T C C T C T C C
Consensus	C C T C C T T T C T	C T C C C C C C	C T C C T C T C C
	940	950	960
seq1	T C T T C T T T C	C T C C C C C C	C C C C T C T C
seq25	T C T T C T T T C	C T C C C C C C	C C C C T C T C
Consensus	T C T T C T T T C	C T C C C C C C	C C C C T C T C
	970	980	990
seq1	T T C T C T C	C C C C C C C	C C C C C C T C
seq25	T T C T C T C	C C C C C C C	C C C C C C T C
Consensus	T T C T C T C	C C C C C C C	C C C C C C T C
	1000	1010	1020
seq1	T C T C T C	C C T C T C T C	T C C C T C C C
seq25	T C T C T C	C C T C T C T C	T C C C T C C C
Consensus	T C T C T C	C C T C T C T C	T C C C T C C C
	1030	1040	1050
seq1	C C C C C	C C C C C	C C T C C C C
seq25	C C C C C	C C C C C	C C T C C C C
Consensus	C C C C C	C C C C C	C C T C C C C



Alignment View Print

	1060	1070	1080
seq1	C G T C T T G	G C C C T C C G	G C C C C T C C G
seq25	C G T C T T G	G C C C T C C G	G C C C C T C C G
Consensus	C G T C T T G	G C C C T C C G	G C C C C T C C G
	1090	1100	1110
seq1	C G T C C C C	G G T G G C C C T G	G G G G C C
seq25	C G T C C C C	G G T G G C C C T G	G G G G C C
Consensus	C G T C C C C	G G T G G C C C T G	G G G G C C
	1120	1130	1140
seq1	T C C C T C T T	C C C G G G C C G	G C C C C C
seq25	T C C C T C T T	C C C G G G C C G	G C C C C C
Consensus	T C C C T C T T	C C C G G G C C G	G C C C C C
	1150	1160	1170
seq1	C T G C C C C	G C T C T G C	C C C C G C G C C
seq25	C T G C C C C	G C T C T G C	C C C C G C G C C
Consensus	C T G C C C C	G C T C T G C	C C C C G C G C C
	1180	1190	1200
seq1	A G G T C C T	G G G C C C C	C T G C T C G
seq25	A G G T C C T	G G G C C C C	C T G C T C G
Consensus	A G G T C C T	G G G C C C C	C T G C T C G
	1210	1220	1230
seq1	G G C T G T C C	C C T G G C C G	C C C C G G C
seq25	G G C T G T C C	C C T G G C C G	C C C C G G C
Consensus	G G C T G T C C	C C T G G C C G	C C C C G G C
	1240	1250	1260
seq1	C C G G G C T T	G G G G C C T C C G	C C C T G G C C T G
seq25	C C G G G C T T	G G G G C C T C C G	C C C T G G C C T G
Consensus	C C G G G C T T	G G G G C C T C C G	C C C T G G C C T G



Alignment View Print

	1270	1280	1290
seq1	CCTTCCCTCT	CCTCCCTCC	CCCTCCCT
seq25	CCTTCCCTCT	CCTCCCTCC	CCCTCCCT
Consensus	CCTTCCCTCT	CCTCCCTCC	CCCTCCCT
	1300	1310	1320
seq1	GGGCCCCCTC	CTGGGGCTC	GCTGGGGCCG
seq25	GGGCCCCCTC	CTGGGGCTC	GCTGGGGCCG
Consensus	GGGCCCCCTC	CTGGGGCTC	GCTGGGGCCG
	1330	1340	1350
seq1	CCCTCTCC	TTCCCCCTGGC	CCCTCTGGGC
seq25	CCCTCTCC	TTCCCCCTGGC	CCCTCTGGGC
Consensus	CCCTCTCC	TTCCCCCTGGC	CCCTCTGGGC
	1360	1370	1380
seq1	CCTTGGGGCCCT	CTGTTCTTCTGCT	GTTGGCCCTGCT
seq25	CCTTGGGGCCCT	CTGTTCTTCTGCT	GTTGGCCCTGCT
Consensus	CCTTGGGGCCCT	CTGTTCTTCTGCT	GTTGGCCCTGCT
	1390	1400	1410
seq1	CTCCCCCGG	TCCTTGGGGAG	GCTGGCCCTGCT
seq25	CTCCCCCGG	TCCTTGGGGAG	GCTGGCCCTGCT
Consensus	CTCCCCCGG	TCCTTGGGGAG	GCTGGCCCTGCT
	1420	1430	1440
seq1	TGCCCCCTTCC	TCCTTGGGGAG	CCCTGGCCCTGCT
seq25	TGCCCCCTTCC	TCCTTGGGGAG	CCCTGGCCCTGCT
Consensus	TGCCCCCTTCC	TCCTTGGGGAG	CCCTGGCCCTGCT
	1450	1460	1470
seq1	CCTCCTTGGCC	CCCTGGGGAG	GGCCCTTCTGCT
seq25	CCTCCTTGGCC	CCCTGGGGAG	GGCCCTTCTGCT
Consensus	CCTCCTTGGCC	CCCTGGGGAG	GGCCCTTCTGCT

Alignment View Print

seq1148014901500
seq25	CCCCCCTC	TCTCCCGCTC	TGCTCTCTCT
Consensus
seq1151015201530
seq25	CCCTCC	CCCGCCC	CGGCGCTGC
Consensus
seq1154015501560
seq25	CTCCGCC	AGGCTGCT	GGCTTGCTG
Consensus
seq1157015801590
seq25	CTCTCTCT	TTTCTGGT	CCCTCTCG
Consensus
seq1160016101620
seq25	GTGATTTT	CTGATTTG	TTTCTTTCC
Consensus
seq1163016401650
seq25	GTCTCTCT	TCTTTTCTC	TCCTCTCT
Consensus
seq1166016701680
seq25	GTCTTTGGGG	GTCTCTCTC	GGGCTCTCT
Consensus

Alignment View Print

seq1 1690 1700 1710
seq25 T T C T T C T T T C C T C
Consensus
seq1 1720 1730 1740
seq25
Consensus

